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TECH CENTER 1600/2900

## SEQUENCE LISTING

<110> Winchester, Robert J.  
Gulko, Percio  
Seki, Tetsunori

<120> USES OF INHIBITORS FOR THE ACTIVATION OF CXCR4  
RECEPTOR BY SDF-1 IN TREATING RHEVMATOID ARTHRITIS

<130> 0575/57005-B

<140> 09/500,746

<141> 2000-02-09

<160> 23

<170> PatentIn version 3.1

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<221> misc\_feature

<223> primer

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gatccgttca tg

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gatcctccct cg

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<400> 7  
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<400> 8  
 gatcctcggt ga

12

<210> 9  
 <211> 507  
 <212> PRT  
 <213> mouse

<220>  
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 <222> (337)..(337)  
 <223> x= to any amino acid

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 <222> (376)..(376)  
 <223> x= to any amino acid

<400> 9

Ser Ala Val Cys Val Tyr His Leu Ser Asp Ile Gln Thr Val Phe Asn  
 1 5 10 15

Gly Pro Phe Ala His Lys Glu Gly Pro Asn His Gln Leu Ile Ser Tyr  
 20 25 30

Gln Gly Arg Ile Pro Tyr Pro Arg Ser Ala Val Cys Val Tyr His Leu  
 35 40 45

Ser Asp Ile Gln Thr Val Phe Asn Gly Pro Phe Ala His Lys Glu Gly

50	55	60
Pro Asn His Gln Leu Ile Ser Tyr Gln Gly Arg Ile Pro Tyr Pro Arg		
65	70	75 80
Ser Ala Val Cys Val Tyr Ser Met Ala Asp Ile Arg Met Val Phe Asn		
	85	90 95
Gly Pro Phe Ala His Lys Glu Gly Pro Asn Tyr Gln Trp Met Pro Phe		
	100	105 110
Ser Gly Lys Met Pro Tyr Pro Arg Ser Ala Val Cys Val Tyr Ser Met		
	115	120 125
Asn Asp Val Arg Arg Ala Phe Leu Gly Pro Phe Ala His Lys Glu Gly		
	130	135 140
Pro Met His Gln Trp Val Ser Tyr Gln Gly Arg Val Pro Tyr Pro Arg		
	145	150 155 160
Ser Ala Val Cys Met Tyr Ser Met Ser Asp Val Arg Arg Val Arg Arg		
	165	170 175
Val Phe Leu Gly Pro Tyr Ala His Arg Asp Gly Pro Asn Tyr Gln Trp		
	180	185 190
Val Pro Tyr Gln Gly Arg Val Pro Tyr Pro Arg Pro Gly Thr Cys Pro		
	195	200 205
Gly Gly Ala Phe Thr Pro Asn Met Arg Thr Thr Lys Asp Phe Pro Asp		
	210	215 220
Asp Val Val Thr Phe Ile Arg Asn His Pro Leu Met Tyr Asn Ser Ile		
	225	230 235 240
Ser Pro Ile Pro Gly Thr Cys Pro Gly Gly Ala Leu Thr Pro Asn Met		
	245	250 255
Arg Thr Thr Lys Glu Phe Pro Asp Asp Val Val Thr Phe Ile Arg Asn		
	260	265 270
His Pro Leu Met Tyr Asn Ser Ile Tyr Pro Ile Pro Gly Thr Cys Pro		
	275	280 285
Gly Gly Thr Phe Thr Pro Ser Met Lys Ser Thr Lys Asp Tyr Pro Asp		
	290	295 300

Glu Val Ile Asn Phe Met Arg Ser His Pro Leu Met Tyr Gln Ala Val  
 305 310 315 320  
 Tyr Pro Leu Pro Gly Met Cys Pro Ser Lys Thr Phe Gly Thr Phe Ser  
 325 330 335  
 Xaa Ser Thr Lys Asp Phe Pro Asp Asp Val Ile Phe Ala Arg Asn His  
 340 345 350  
 Pro Leu Met Tyr Asn Ser Val Leu Pro Thr Pro Gly Thr Cys Pro Ser  
 355 360 365  
 Lys Thr Phe Gly Gly Phe Asp Xaa Ser Thr Lys Asp Leu Pro Asp Asp  
 370 375 380  
 Val Ile Thr Phe Ala Arg Ser His Pro Ala Met Tyr Asn Pro Val Phe  
 385 390 395 400  
 Pro Met His Arg Arg Pro Leu Ile Val Arg Ile Gly Thr Asp Tyr Lys  
 405 410 415  
 Tyr Thr Lys Ile Ala Val Asp His Lys Arg Pro Leu Ile Val Arg Ile  
 420 425 430  
 Gly Thr Asp Tyr Lys Tyr Thr Lys Ile Ala Val Asp Gln Arg Arg Pro  
 435 440 445  
 Leu Val Val Arg Thr Gly Ala Pro Tyr Arg Leu Thr Thr Ile Ala Val  
 450 455 460  
 Asp Gly Gly Arg Pro Leu Phe Leu Gln Val Gly Ala Asn Tyr Thr Phe  
 465 470 475 480  
 Thr Gln Ile Ala Ala Asp Asn Asn Arg Pro Ile Val Ile Lys Thr Asp  
 485 490 495  
 Val Asn Tyr Gln Phe Thr Gln Ile Val Val Asp  
 500 505

<210> 10  
 <211> 396  
 <212> PRT  
 <213> Human

<400> 10

Ser Tyr Pro Ala Pro His Gly Pro Glu Asp Pro Ala Pro Gln Phe Ala  
 1 5 10 15

His Met Phe Glu Asn Glu Ile Ser His Arg Thr Gly Ser Trp Asn Phe  
 20 25 30  
 Ala Pro Asn Pro Asp Lys Gln Trp Leu Leu Gln Arg Thr Ser His Ala  
 35 40 45  
 Ala Pro His Gly Pro Glu Asp Ser Ala Pro Gln Phe Ser Glu Leu Tyr  
 50 55 60  
 Pro Asn Ala Ser Gln His Ile Thr Pro Ser Tyr Asn Tyr Ala Pro Asn  
 65 70 75 80  
 Met Asp Lys His Trp Ile Met Gln Tyr Thr Ala Thr Pro Ala Pro His  
 85 90 95  
 Ser Pro Trp Thr Ala Ala Pro Gln Tyr Gln Lys Ala Phe Gln Asn Val  
 100 105 110  
 Phe Ala Pro Arg Asn Lys Asn Phe Asn Ile His Gly Thr Asn Lys His  
 115 120 125  
 Trp Leu Ile Arg Gln Ala Lys Gly Lys Met Asn Asp Val His Ile Ser  
 130 135 140  
 Phe Thr Asp Leu Leu His Arg Arg Arg Leu Gln Thr Leu Gln Ser Val  
 145 150 155 160  
 Asp Glu Gly Ile Glu Arg Leu Phe Asn Leu Leu Arg Glu Leu Asn Gln  
 165 170 175  
 Leu Trp Asn Thr Gly Pro Met Leu Pro Ile His Met Glu Phe Thr Asn  
 180 185 190  
 Ile Leu Gln Arg Lys Arg Leu Gln Thr Leu Met Ser Val Asp Asp Ser  
 195 200 205  
 Val Glu Arg Leu Tyr Asn Met Leu Val Glu Thr Gly Glu Leu Glu Asn  
 210 215 220  
 Thr Thr Pro Met Thr Asn Ser Ser Ile Gln Phe Leu Asp Asn Ala Phe  
 225 230 235 240  
 Arg Lys Arg Trp Gln Thr Leu Leu Ser Val Asp Asp Leu Val Glu Lys  
 245 250 255

Leu Val Lys Arg Leu Glu Phe Thr Gly Glu Leu Asn Asn Thr Tyr Ala  
 260 265 270

Ile Tyr Thr Ser Asp His Gly Tyr His Leu Gly Gln Phe Gly Leu Leu  
 275 280 285

Lys Gly Lys Asn Met Pro Tyr Glu Phe Asp Ile Arg Val Pro Phe Phe  
 290 295 300

Met Arg Gly Pro Gly Ile Pro Arg Tyr Ile Ile Tyr Thr Ala Asp His  
 305 310 315 320

Gly Tyr His Ile Gly Gln Phe Gly Leu Val Lys Gly Lys Ser Met Pro  
 325 330 335

Tyr Asp Phe Asp Ile Arg Val Pro Phe Phe Ile Arg Gly Pro Ser Val  
 340 345 350

Glu Pro Tyr Ile Phe Tyr Thr Ser Asp Asn Gly Tyr His Thr Gly Gln  
 355 360 365

Phe Ser Leu Pro Ile Asp Lys Arg Gln Leu Tyr Glu Phe Asp Ile Lys  
 370 375 380

Val Pro Leu Leu Val Arg Gly Pro Gly Ile Lys Pro  
 385 390 395

<210> 11  
 <211> 102  
 <212> PRT  
 <213> Human

<400> 11

Ser Ala Val Cys Val Tyr Tyr Ser Met Ala Asp Ile Arg Met Val Phe  
 1 5 10 15

Asn Gly Pro Phe Ala His Lys Glu Gly Pro Asn Tyr Gln Trp Met Pro  
 20 25 30

Phe Ser Gly Lys Met Pro Tyr Pro Arg Pro Gly Thr Cys Pro Gly Gly  
 35 40 45

Thr Phe Thr Pro Ser Met Lys Ser Thr Lys Asx Tyr Pro Asp Glu Val  
 50 55 60

Ile Asn Phe Met Arg Ser His Pro Leu Met Tyr Gln Ala Val Tyr Pro  
 65 70 75 80

Leu Gln Arg Arg Pro Leu Val Val Arg Thr Gly Ala Pro Tyr Arg Leu  
                           85                          90                          95

Thr Thr Ile Ala Val Asp  
                           100

<210> 12  
 <211> 101  
 <212> PRT  
 <213> Human

<220>  
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 <222> (54)..(54)  
 <223> X= to any amino acid

<400> 12

Ser Ala Val Cys Val Tyr Ser Met Asn Asp Val Arg Arg Ala Phe Leu  
   1                          5                          10                          15

Gly Pro Phe Ala His Lys Glu Gly Pro Met His Gln Trp Val Ser Tyr  
                           20                          25                          30

Gln Gly Arg Val Pro Tyr Pro Arg Pro Gly Met Cys Pro Ser Lys Thr  
                           35                          40                          45

Phe Gly Thr Phe Ser Xaa Ser Thr Lys Asp Phe Pro Asp Asp Val Ile  
   50                          55                          60

Gln Phe Ala Arg Asn His Pro Lys Met Tyr Asn Ser Val Leu Pro Thr  
   65                          70                          75                          80

Gly Gly Arg Pro Leu Phe Leu Gln Val Gly Ala Asn Tyr Thr Phe Thr  
                           85                          90                          95

Gln Ile Ala Ala Asp  
                           100

<210> 13  
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 <212> PRT  
 <213> Human

<220>  
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 <222> (54)..(54)  
 <223> X=to any amino acid

<400> 13



Ser Ala Val Cys Met Tyr Ser Met Ser Asp Val Arg Arg Val Phe Leu  
1 5 10 15

Gly Pro Tyr Ala His Arg Asp Gly Pro Asn Tyr Gln Trp Val Pro Tyr  
20 25 30

Gln Gly Arg Val Pro Tyr Pro Arg Pro Gly Thr Cys Pro Ser Lys Thr  
35 40 45

Phe Gly Gly Phe Asp Xaa Ser Thr Lys Asp Leu Pro Asp Asp Val Ile  
50 55 60

Thr Phe Ala Arg Ser His Pro Ala Met Tyr Asn Pro Val Phe Pro Met  
65 70 75 80

Asn Asn Arg Pro Ile Val Ile Lys Thr Asp Val Asn Tyr Gln Phe Thr  
85 90 95

Gln Ile Val Val Asp  
100

<210> 14  
<211> 90  
<212> PRT  
<213> worm

<400> 14

Ser Tyr Pro Ala Pro His Gly Pro Glu Asp Pro Ala Pro Gln Phe Ala  
1 5 10 15

His Met Phe Glu Asn Glu Ile Ser His Arg Thr Gly Ser Trp Asn Phe  
20 25 30

Ala Pro Asn Pro Asp Lys Gln Trp Leu Leu Gln Arg Thr Gly Lys Met  
35 40 45

Asn Asp Val His Ile Ser Phe Thr Asp Leu Leu His Arg Arg Arg Leu  
50 55 60

Gln Thr Leu Gln Ser Val Asp Glu Gly Ile Glu Arg Leu Phe Asn Leu  
65 70 75 80

Leu Arg Glu Leu Asn Gln Leu Trp Asn Thr  
85 90

<210> 15  
<211> 132

<212> PRT  
<213> Worm

<400> 15

Ser His Ala Ala Pro His Gly Pro Glu Asp Ser Ala Pro Gln Phe Ser  
1 5 10 15

Glu Leu Tyr Pro Asn Ala Ser Gln His Ile Thr Pro Ser Tyr Asn Tyr  
20 25 30

Ala Pro Asn Met Asp Lys His Trp Ile Met Gln Tyr Thr Gly Pro Met  
35 40 45

Leu Pro Ile His Met Glu Phe Thr Asn Ile Leu Gln Arg Lys Arg Leu  
50 55 60

Gln Thr Leu Met Ser Val Asp Asp Ser Val Glu Arg Leu Tyr Asn Met  
65 70 75 80

Leu Val Glu Thr Gly Glu Leu Glu Asn Thr Tyr Ile Ile Tyr Thr Ala  
85 90 95

Asp His Gly Tyr His Ile Gly Gln Phe Gly Leu Val Lys Gly Lys Ser  
100 105 110

Met Pro Tyr Asp Phe Asp Ile Arg Val Pro Phe Phe Ile Arg Gly Pro  
115 120 125

Ser Val Glu Pro  
130

<210> 16  
<211> 130  
<212> PRT  
<213> Human

<400> 16

Ala Thr Pro Ala Pro His Ser Pro Trp Thr Ala Ala Pro Gln Lys Ala  
1 5 10 15

Phe Gln Asn Val Phe Ala Pro Arg Asn Lys Asn Phe Asn Ile His Gly  
20 25 30

Thr Asn Lys His Trp Leu Ile Arg Gln Ala Lys Thr Pro Met Thr Asn  
35 40 45

Ser Ser Ile Gln Phe Leu Asp Asn Ala Phe Arg Lys Arg Trp Gln Thr  
50 55 60

Leu Leu Ser val Asp Asp Leu val Glu Lys Leu val Lys Arg Leu Glu  
65 70 75 80

Phe Thr Gly Glu Leu Asn Asn Thr Tyr Ile Phe Tyr Thr Ser Asp Asn  
85 90 95

Gly Tyr His Thr Gly Gln Phe Ser Leu Pro Ile Asp Lys Arg Gln Leu  
100 105 110

Tyr Glu Phe Asp Ile Lys val Pro Leu Leu val Arg Gly Pro Gly Ile  
115 120 125

Lys Pro  
130

<210> 17  
<211> 410  
<212> PRT  
<213> Human

<400> 17

Gly Asn Asn Gly Ala Gly Thr Gly Thr Gly Gly Gly Ala Cys Gly Gly  
1 5 10 15

Gly Gly Gly Gly Asn Gly Asn Ala Gly Asn Ala Ala Thr Thr Ala Ala  
20 25 30

Gly Gly Thr Ala Gly Asn Gly Ala Thr Gly Gly Ala Gly Asn Ala Asn  
35 40 45

Gly Gly Gly Gly Thr Gly Cys Asn Thr Asn Gly Gly Asn Asn Asn Ala  
50 55 60

Gly Ala Asn Ala Asn Thr Gly Asn Asn Thr Gly Gly Ala Gly Ala Ala  
65 70 75 80

Asn Gly Ala Cys Ala Ala Asn Gly Gly Gly Gly Gly Asn Gly Thr Cys  
85 90 95

Gly Asn Asn Gly Gly Ala Gly Cys Asn Gly Asn Thr Gly Thr Gly Ala  
100 105 110

Gly Thr Gly Gly Gly Ala Ala Gly Ala Ala Gly Gly Cys Asn Ala Cys  
115 120 125

Gly Thr Cys Ala Ala Asn Ala Ala Gly Gly Ala Cys Gly Ala Ala Thr

130

135

140

Ala Thr Thr Thr Gly Cys Ala Ala Asn Gly Asn Asn Gly Asn Asn Cys  
 145 150 155 160

Ala Gly Gly Gly Cys Thr Gly Thr Asn Cys Asn Cys Gly Gly Gly Cys  
 165 170 175

Ala Gly Thr Thr Thr Gly Thr Ala Ala Ala Ala Ala Ala Ala Ala Ala  
 180 185 190

Ala Ala Asn Ala Ala Gly Ala Ala Cys Asn Gly Cys Gly Ala Cys Ala  
 195 200 205

Gly Ala Cys Ala Ala Gly Thr Gly Thr Asn Asn Gly Thr Thr Gly Ala  
 210 215 220

Cys Cys Cys Gly Ala Ala Gly Cys Asn Ala Asn Ala Gly Thr Gly Gly  
 225 230 235 240

Ala Thr Asn Cys Ala Gly Gly Ala Gly Thr Ala Cys Cys Thr Gly Gly  
 245 250 255

Ala Gly Asn Asn Ala Ala Cys Thr Ala Thr Gly Ala Ala Cys Ala Ala  
 260 265 270

Asn Thr Ala Ala Gly Cys Gly Cys Ala Ala Cys Ala Gly Cys Cys Ala  
 275 280 285

Ala Ala Gly Ala Gly Gly Ala Cys Thr Thr Asn Cys Cys Gly Cys Thr  
 290 295 300

Ala Gly Ala Cys Cys Cys Ala Cys Thr Cys Gly Ala Gly Gly Ala Ala  
 305 310 315 320

Ala Ala Cys Thr Ala Ala Ala Ala Cys Cys Thr Thr Gly Thr Gly Ala  
 325 330 335

Gly Ala Gly Ala Thr Gly Ala Ala Ala Gly Gly Asn Cys Ala Ala Ala  
 340 345 350

Gly Ala Cys Gly Thr Gly Gly Gly Gly Gly Ala Gly Gly Gly Gly Gly  
 355 360 365

Cys Cys Asn Thr Ala Ala Cys Cys Ala Thr Gly Ala Gly Gly Ala Cys  
 370 375 380

Cys Ala Gly Gly Thr Gly Thr Gly Thr Gly Thr Gly Thr Gly Thr Gly  
 385 390 395 400

Thr Gly Gly Gly Gly Thr Gly Gly Gly Cys  
 405 410

<210> 18  
 <211> 425  
 <212> PRT  
 <213> Human

<400> 18

Cys Cys Cys Gly Gly Gly Thr Ala Cys Cys Gly Ala Gly Cys Thr Cys  
 1 5 10 15

Gly Ala Ala Thr Thr Cys Cys Gly Thr Thr Gly Asn Thr Gly Thr Cys  
 20 25 30

Gly Cys Cys Gly Thr Thr Gly Asn Thr Gly Thr Cys Gly Cys Ala Gly  
 35 40 45

Ala Thr Gly Cys Cys Cys Ala Thr Gly Cys Cys Cys Ala Thr Gly Cys  
 50 55 60

Cys Gly Ala Thr Thr Cys Thr Thr Cys Gly Ala Ala Ala Gly Cys Cys  
 65 70 75 80

Ala Thr Gly Thr Thr Gly Cys Cys Ala Gly Ala Gly Cys Cys Ala Ala  
 85 90 95

Cys Gly Thr Cys Ala Ala Gly Cys Ala Thr Cys Thr Cys Ala Ala Ala  
 100 105 110

Ala Thr Thr Cys Thr Cys Ala Ala Cys Ala Cys Thr Cys Cys Ala Ala  
 115 120 125

Ala Cys Thr Gly Thr Gly Cys Cys Cys Thr Thr Cys Ala Gly Ala Thr  
 130 135 140

Thr Gly Thr Ala Gly Cys Cys Cys Gly Gly Cys Thr Gly Ala Ala Gly  
 145 150 155 160

Ala Ala Cys Ala Ala Cys Ala Ala Cys Ala Gly Ala Cys Ala Ala Gly  
 165 170 175

Thr Gly Thr Gly Cys Ala Thr Thr Gly Ala Cys Cys Cys Gly Ala Ala  
 180 185 190

Gly Cys Thr Ala Ala Ala Gly Thr Gly Gly Ala Thr Thr Cys Ala Gly  
 195 200 205  
 Gly Ala Gly Thr Ala Cys Cys Thr Gly Gly Ala Gly Ala Ala Ala Gly  
 210 215 220  
 Cys Thr Thr Thr Ala Ala Ala Cys Ala Ala Gly Thr Ala Ala Gly Cys  
 225 230 235 240  
 Ala Cys Ala Ala Cys Ala Gly Cys Cys Ala Ala Ala Ala Ala Gly Gly  
 245 250 255  
 Ala Cys Thr Thr Thr Cys Cys Gly Cys Thr Ala Gly Ala Cys Cys Cys  
 260 265 270  
 Ala Asn Thr Cys Gly Ala Gly Ala Ala Ala Ala Cys Thr Ala Ala Ala  
 275 280 285  
 Ala Cys Cys Thr Thr Gly Thr Gly Ala Gly Ala Gly Ala Thr Gly Ala  
 290 295 300  
 Ala Ala Gly Gly Gly Cys Ala Ala Ala Gly Ala Cys Gly Thr Gly Gly  
 305 310 315 320  
 Gly Gly Gly Gly Ala Gly Gly Gly Gly Gly Gly Cys Thr Thr Ala Ala  
 325 330 335  
 Cys Cys Ala Thr Gly Ala Gly Gly Ala Cys Cys Ala Gly Gly Thr Gly  
 340 345 350  
 Thr Gly Thr Gly Thr Gly Thr Asn Gly Gly Gly Thr Gly Gly Gly Gly  
 355 360 365  
 Cys Ala Cys Ala Thr Thr Gly Gly Ala Thr Cys Thr Thr Asn Gly Ala  
 370 375 380  
 Thr Cys Gly Gly Gly Cys Cys Thr Gly Ala Gly Gly Thr Thr Thr Gly  
 385 390 395 400  
 Gly Cys Ala Gly Cys Ala Thr Thr Thr Ala Gly Ala Cys Cys Cys Thr  
 405 410 415  
 Gly Gly Ala Thr Thr Ala Thr Gly Asn  
 420 425

<210> 19  
 <211> 376  
 <212> PRT  
 <213> Human

<400> 19

Cys Ala Gly Ala Thr Gly Cys Cys Cys Ala Thr Gly Cys Cys Gly Ala  
 1 5 10 15

Thr Thr Cys Thr Thr Cys Gly Ala Ala Gly Cys Cys Ala Thr Gly  
 20 25 30

Thr Thr Gly Cys Cys Ala Gly Ala Gly Cys Cys Ala Ala Cys Gly Thr  
 35 40 45

Cys Ala Ala Gly Cys Ala Thr Cys Thr Cys Ala Ala Ala Thr Thr  
 50 55 60

Cys Thr Cys Ala Ala Cys Ala Cys Thr Cys Cys Ala Ala Ala Cys Thr  
 65 70 75 80

Gly Thr Gly Cys Cys Cys Thr Thr Cys Ala Gly Ala Thr Thr Gly Thr  
 85 90 95

Ala Gly Cys Cys Cys Gly Gly Cys Thr Gly Ala Ala Gly Ala Ala Cys  
 100 105 110

Ala Ala Cys Ala Ala Cys Ala Gly Ala Cys Ala Ala Gly Thr Gly Thr  
 115 120 125

Gly Cys Ala Thr Thr Gly Ala Cys Cys Cys Gly Ala Ala Gly Cys Thr  
 130 135 140

Ala Ala Ala Gly Thr Gly Gly Ala Thr Thr Cys Ala Gly Gly Ala Gly  
 145 150 155 160

Thr Ala Cys Cys Thr Gly Gly Ala Gly Gly Ala Ala Ala Gly Cys Thr  
 165 170 175

Thr Thr Ala Ala Ala Cys Ala Ala Gly Thr Ala Ala Gly Cys Ala Cys  
 180 185 190

Ala Ala Cys Ala Gly Cys Cys Ala Ala Ala Ala Ala Gly Gly Ala Cys  
 195 200 205

Thr Thr Thr Cys Cys Gly Cys Thr Ala Gly Ala Cys Cys Cys Ala Cys  
 210 215 220

Thr Cys Gly Ala Gly Gly Ala Ala Ala Ala Cys Thr Ala Ala Ala Ala  
225 230 235 240

Cys Cys Thr Thr Gly Thr Gly Ala Gly Ala Gly Ala Thr Gly Ala Ala  
245 250 255

Ala Gly Gly Gly Cys Ala Ala Asn Gly Ala Cys Gly Thr Asn Gly Asn  
260 265 270

Gly Gly Ala Gly Gly Gly Gly Gly Gly Cys Thr Thr Ala Ala Cys Cys  
275 280 285

Ala Thr Gly Ala Gly Gly Ala Cys Cys Ala Gly Gly Thr Gly Thr Gly  
290 295 300

Thr Asn Thr Gly Gly Gly Gly Gly Thr Gly Gly Gly Thr Ala Cys Ala  
305 310 315 320

Thr Thr Gly Asn Ala Thr Cys Thr Thr Gly Gly Gly Ala Thr Cys Gly  
325 330 335

Gly Gly Cys Cys Thr Gly Ala Gly Gly Thr Thr Asn Gly Gly Cys Ala  
340 345 350

Gly Ala Ala Thr Thr Thr Asn Gly Asn Cys Cys Cys Thr Gly Asn Ala  
355 360 365

Thr Thr Thr Asn Thr Gly Gly Asn  
370 375

<210> 20  
<211> 377  
<212> PRT  
<213> Human

<400> 20

Cys Ala Gly Ala Thr Gly Asn Cys Cys Ala Thr Gly Cys Cys Gly Ala  
1 5 10 15

Thr Thr Cys Thr Thr Cys Gly Ala Ala Ala Gly Cys Cys Ala Thr Gly  
20 25 30

Thr Thr Gly Cys Cys Ala Gly Ala Gly Cys Cys Ala Ala Cys Gly Thr  
35 40 45

Cys Ala Ala Gly Cys Ala Thr Cys Thr Cys Ala Ala Ala Ala Thr Thr  
50 55 60



Cys Thr Cys Ala Ala Cys Ala Cys Thr Cys Cys Ala Ala Ala Cys Thr  
 65 70 75 80  
 Gly Thr Gly Cys Cys Cys Thr Thr Cys Ala Gly Ala Thr Thr Gly Thr  
 85 90 95  
 Ala Gly Cys Cys Cys Gly Gly Cys Thr Gly Ala Ala Gly Ala Ala Cys  
 100 105 110  
 Ala Ala Cys Ala Ala Cys Ala Gly Ala Cys Ala Ala Gly Thr Gly Thr  
 115 120 125  
 Gly Cys Ala Thr Thr Gly Ala Cys Cys Cys Gly Ala Ala Gly Cys Thr  
 130 135 140  
 Ala Ala Ala Gly Thr Gly Gly Ala Thr Thr Cys Ala Gly Gly Ala Gly  
 145 150 155 160  
 Thr Ala Cys Cys Thr Gly Gly Ala Gly Thr Ala Ala Ala Gly Cys Thr  
 165 170 175  
 Thr Thr Ala Ala Ala Cys Ala Ala Gly Thr Ala Ala Gly Cys Ala Cys  
 180 185 190  
 Ala Ala Cys Ala Gly Asn Cys Ala Ala Ala Ala Ala Gly Gly Ala Cys  
 195 200 205  
 Thr Thr Thr Cys Cys Gly Cys Thr Ala Gly Ala Cys Cys Cys Ala Cys  
 210 215 220  
 Thr Cys Gly Ala Gly Gly Ala Ala Ala Ala Cys Thr Ala Ala Ala Ala  
 225 230 235 240  
 Cys Cys Thr Thr Gly Thr Gly Ala Gly Ala Gly Ala Thr Gly Ala Ala  
 245 250 255  
 Ala Gly Gly Gly Cys Ala Asn Thr Gly Thr Thr Asn Thr Thr Gly Thr  
 260 265 270  
 Gly Gly Ala Gly Gly Gly Gly Gly Cys Cys Thr Thr Ala Ala Cys Cys  
 275 280 285  
 Ala Thr Gly Ala Gly Gly Ala Cys Cys Ala Gly Gly Thr Gly Thr Gly  
 290 295 300

Thr Gly Thr Gly Thr Gly Gly Gly Gly Thr Gly Gly Gly Cys Ala Cys  
305 310 315 320

Ala Thr Asn Gly Asn Ala Thr Cys Thr Gly Gly Gly Thr Ala Thr Cys  
325 330 335

Gly Gly Gly Cys Cys Thr Gly Ala Gly Gly Thr Thr Thr Gly Asn Cys  
340 345 350

Ala Gly Cys Ala Thr Thr Thr Ala Gly Asn Cys Cys Cys Thr Gly Asn  
355 360 365

Ala Thr Thr Thr Ala Thr Asn Gly Cys  
370 375

<210> 21  
<211> 292  
<212> PRT  
<213> Human

<400> 21

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1 5 10 15

Ala Cys Gly Thr Cys Ala Ala Cys Ala Thr Cys Cys Cys Ala Ala Ala  
20 25 30

Ala Thr Cys Thr Cys Ala Ala Cys Ala Cys Asn Cys Cys Cys Ala Ala  
35 40 45

Cys Thr Asn Thr Thr Cys Cys Cys Thr Thr Cys Ala Gly Ala Thr Thr  
50 55 60

Gly Thr Ala Gly Cys Cys Cys Gly Gly Cys Thr Gly Ala Ala Gly Ala  
65 70 75 80

Ala Cys Ala Ala Cys Ala Ala Cys Ala Ala Gly Ala Cys Ala Ala Gly  
85 90 95

Thr Gly Thr Gly Cys Ala Thr Thr Thr Gly Ala Cys Cys Cys Gly Ala  
100 105 110

Ala Gly Cys Thr Ala Ala Ala Ala Gly Thr Gly Gly Ala Thr Thr Cys  
115 120 125

Ala Gly Gly Ala Gly Thr Ala Cys Cys Cys Thr Gly Gly Ala Gly Ala  
130 135 140

Ala Ala Gly Cys Thr Thr Thr Ala Ala Ala Cys Ala Ala Gly Thr Ala  
145 150 155 160

Ala Gly Cys Ala Cys Ala Ala Cys Ala Gly Cys Cys Cys Ala Ala Ala  
165 170 175

Ala Ala Gly Gly Ala Cys Thr Thr Thr Cys Cys Gly Cys Thr Ala Gly  
180 185 190

Ala Cys Cys Cys Ala Cys Thr Cys Gly Ala Gly Gly Ala Ala Ala Ala  
195 200 205

Cys Thr Ala Ala Ala Ala Cys Cys Thr Thr Gly Thr Gly Ala Gly Ala  
210 215 220

Gly Ala Thr Gly Ala Ala Ala Gly Gly Asn Cys Ala Ala Ala Gly Ala  
225 230 235 240

Cys Gly Thr Gly Gly Gly Gly Gly Ala Gly Gly Gly Gly Gly Cys Cys  
245 250 255

Thr Thr Ala Ala Cys Cys Ala Thr Gly Ala Gly Gly Ala Cys Cys Ala  
260 265 270

Gly Gly Thr Gly Thr Gly Thr Gly Thr Gly Thr Gly Gly Gly Thr  
275 280 285

Gly Gly Gly Cys  
290

<210> 22  
<211> 75  
<212> PRT  
<213> Human

<400> 22

Ala Asn Thr Gly Ala Ala Gly Gly Gly Cys Cys Ala Ala Ala Gly Ala  
1 5 10 15

Cys Gly Thr Gly Gly Gly Gly Gly Ala Gly Gly Gly Gly Gly Cys Cys  
20 25 30

Thr Thr Ala Ala Cys Cys Cys Ala Thr Thr Gly Ala Gly Gly Ala Cys  
35 40 45

Cys Ala Gly Asn Thr Gly Thr Gly Thr Gly Thr Gly Gly Gly Gly Gly  
50 55 60

Thr Gly Gly Gly Gly Gly Thr Gly Gly Cys Cys  
 65 70 75

<210> 23  
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<220>  
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 <222> (2)..(2)  
 <223> X = to any amino acid

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 <223> X = to any amino acid

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 <223> X = to any amino acid

<400> 23

Gly Xaa Xaa Gly Ala Gly Thr Gly Thr Gly Gly Gly Ala Cys Gly Gly

1	5	10	15
Gly Gly Gly Xaa Gly Xaa Ala Ala Thr Thr Ala Ala Gly Ser Tyr Met	20	25	30
Gly Gly Gly Thr Ala Tyr Ser Gly Ala Gly Cys Trp Cys Gly Arg Arg	35	40	45
Lys Thr Ser Cys Gly Thr Thr Gly Gly Thr Gly Thr Met Gly Met Cys	50	55	60
Arg Thr Thr Gly Xaa Xaa Thr Gly Lys Met Gly Ala Ala Xaa Gly Ala	65	70	80
Cys Ala Gly Ala Thr Gly Ser Cys Cys Ala Thr Gly Cys Cys Gly Ala	85	90	95
Thr Thr Cys Thr Thr Cys Gly Ala Ala Ala Gly Cys Cys Ala Thr Gly	100	105	110
Thr Thr Gly Cys Met Ala Gly Ala Gly Cys Cys Ala Ala Cys Gly Thr	115	120	125
Cys Ala Ala Gly Cys Ala Thr Cys His Cys Ala Ala Ala Ala Thr Thr	130	135	140
Cys Thr Cys Ala Ala Cys Ala Cys Thr Cys Cys Met Ala Ala Cys Thr	145	150	155
Gly Thr Gly Cys Cys Cys Thr Thr Cys Ala Gly Ala Thr Thr Gly Thr	165	170	175
Ala Gly Cys Cys Cys Gly Gly Cys Thr Gly Ala Ala Gly Ala Ala Cys	180	185	190
Ala Ala Cys Ala Ala Cys Ala Ala Gly Ala Cys Ala Ala Gly Thr Gly	195	200	205
Thr Gly Thr Gly Cys Ala Thr Thr Gly Ala Cys Cys Cys Gly Ala Ala	210	215	220
Gly Cys Thr Ala Ala Ala Ala Gly Thr Gly Gly Ala Thr Thr Cys Ala	225	230	235
Gly Gly Ala Gly Thr Ala Cys Cys Thr Gly Gly Ala Gly Lys Ala Ala	245	250	255

Ala Gly Cys Thr Thr Thr Ala Ala Ala Cys Ala Ala Gly Thr Ala Ala  
260 265 270

Gly Cys Ala Cys Ala Ala Cys Ala Gly Cys Cys Cys Ala Ala Ala Ala  
275 280 285

Ala Gly Gly Ala Cys Thr Thr Thr Cys Cys Gly Cys Thr Ala Gly Ala  
290 300

Cys Cys Cys Ala Cys Thr Cys Gly Ala Gly Gly Ala Ala Ala Ala Cys  
305 310 315 320

Thr Ala Ala Ala Ala Cys Cys Thr Thr Gly Thr Gly Ala Gly Ala Gly  
325 330 335

Ala Thr Gly Ala Ala Ala Gly Gly Ser Cys Ala Ala Trp Gly Ala Cys  
340 345 350

Gly Thr Lys Gly Lys Gly Gly Ala Gly Gly Gly Gly Gly Ser Cys Thr  
355 360 365

Thr Ala Ala Cys Cys Cys Ala Thr Thr Gly Ala Gly Gly Ala Cys Cys  
370 375 380

Ala Gly Gly Thr Gly Thr Gly Thr Gly Thr Gly Gly Gly Gly Gly Thr  
385 390 395 400

Gly Gly Cys Ala Cys Ala Thr Thr Gly Xaa Ala Thr Cys Thr Thr Gly  
405 410 415

Gly Gly Ala Thr Cys Gly Gly Gly Cys Cys Thr Gly Ala Gly Gly Thr  
420 425 430

Thr Thr Gly Ser Cys Ala Gly Cys Ala Thr Thr Thr Ala Gly Ala Cys  
435 440 445

Cys Cys Thr Gly Ser Ala Thr Thr Thr Ala Thr Arg Gly Cys  
450 455 460